



SEQUENCE LISTING

<110> Takano, Eriko
<111> Eiko, Meryn

<120> Anticanceric Production

<130> 1381-P-0329US1

<140> 11-017,471

<141> 2001-10-23

<150> 60-048,561

<161> 2000-10-23

<160> 19

<170> PatentIn Ver. 2.1

<210> 1

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 1

gaccacgtsc csggcatg

18

<210> 2

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 2

gtccctgctgg ccgtsacsc gsac

24

<210> 3

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 3

gcacatggt ggggacg ac

22

<210> 4

<211> 15

<212> DNA

<213> Artificial Sequence

<210>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide

 <410> 4
 gataccggg aacgcttggc catg 15

 <210> 5
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide

 <410> 5
 tataragctg accgggaacg cgtc 24

 <210> 6
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide

 <410> 6
 atcgcccggt cctgcttggc catg 24

 <210> 7
 <211> 22
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide

 <410> 7
 aagtagaggg ctcccttggt ca 22

 <210> 8
 <211> 22
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Synthetic
 oligonucleotide

 <410> 8
 gaaaactact gtttagggc ag 22

 <210> 9
 <211> 21
 <212> DNA
 <213> Artificial Sequence

<210>
<213> Description of Artificial Sequence: Synthetic
oligonucleotide

<410> 9
ctgcacccctg gtcgggtgga ca

22

<210> 11
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 10
atcgaccgggt cctgcttgga catg

24

<210> 11
<211> 31
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 11
catctgcagc gtgatcgtag cagcttggtg g

31

<210> 12
<211> 401
<212> DNA
<213> Streptomyces coelicolor

<400> 12
gggagaggacg ggggtgacg agaacgggtc accgccccttc ggtatccagc tgaaccgggaa 60
cgggtccttgc accctgggtc ggtggacaag cggcatcgga accgggaatg cggtttggtc 120
gatcgagttg gcatcggtac cagaattgat caaaactact gtttcgggca tgggttcccc 180
ccaggaaatca tgtgatgccc agctgtttctg tatgcgggaa cyttaagata cagactgagc 240
ggtttttttt ctatcccttc cgggggagac atgaacaagg aggcaggcat ggccaagcag 300
gacgggggga tccgcacggg gcagacgata ctggacggcg cggcgaggt ctccgagaag 360
cagygctacc aagctggcac gatcacggag atcctcaagg c 401

<210> 12
<211> 401
<212> DNA
<213> Streptomyces coelicolor

<400> 12
accttgagga tctccgtgat cgtggcagct tggtagccct gcttctcgaa gacctgcgcc 60
gggggttcca ggatggtctg ccgctggcgg atcgcccggt cctgcttgga catgcctgcc 120
tcttggttca tgtctccccc gggaaggata gaaaaaaaaa cgttcagttc gtatottaac 180
gttcgggcat acagaacagc tgggcacac atgattccctg ggggggaccc atgcgccag 240
cagtatgttt gatcaattct ggttcggatg ccaactcgat cgaacaaaac gaattgcggg 300
tccgatggg gcttgtccac cggaccaggg tgcaggacgc gttcccggtt agctgggatac 360
cgaagggggg tgaaccgttc tgggtacgg cgttcctgcc c 401

<210> 14
<211> 67
<212> PRT
<213> Streptomyces coelicolor

<400> 14
Met Pro Glu Ala Val Val Leu Ile Asn Ser Ala Ser Asp Ala Asn Ser
1 5 10 15
Ile Glu Gln Thr Ala Leu Pro Val Pro Met Ala Leu Val His Arg Thr
20 25 30
Arg Val Gln Asp Ala Phe Pro Val Ser Trp Ile Pro Lys Gly Gly Asp
35 40 45
Arg Phe Ser Val Thr Ala Val Leu Pro
50 55

<210> 15
<211> 37
<212> PRT
<213> Streptomyces coelicolor

<400> 15
Met Ala Lys Gln Asp Arg Ala Ile Arg Thr Arg Gln Thr Ile Leu Asp
1 5 10 15
Ala Ala Ala Gln Val Phe Glu Lys Gln Gly Tyr Gln Ala Ala Thr Ile
20 25 30
Thr Glu Ile Leu Lys
35

<210> 16
<211> 215
<212> PRT
<213> Streptomyces coelicolor

<400> 16
Met Ala Lys Gln Asp Arg Ala Ile Arg Thr Arg Gln Thr Ile Leu Asp
1 5 10 15
Ala Ala Ala Gln Val Phe Glu Lys Gln Gly Tyr Gln Ala Ala Thr Ile
20 25 30
Thr Glu Ile Leu Lys Val Ala Gly Val Thr Lys Gly Ala Leu Tyr Phe
35 40 45
His Phe Gln Ser Lys Glu Glu Leu Ala Leu Gly Val Phe Asp Ala Gln
50 55 60
Glu Pro Pro Gln Ala Val Pro Glu Gln Pro Leu Arg Leu Glu Glu Leu
65 70 75 80
Ile Asp Met Gly Met Leu Phe Cys His Arg Leu Arg Thr Asn Val Val
85 90 95
Ala Arg Ala Gly Val Arg Leu Ser Met Asp Glu Glu Ala His Gly Leu
100 105 110
Asp Arg Arg Gly Pro Phe Arg Arg Trp His Glu Thr Leu Leu Lys Leu
115 120 125

Leu Asn Gln Ala Lys Glu Asn Gly Gln Leu Leu Pro His Val Val Thr
 130 135 140
 Thr Asp Ser Ala Asp Leu Tyr Val Gly Thr Phe Ala Gly Ile Gln Val
 145 150 155 160
 Val Ser Gln Thr Val Ser Asp Tyr Gln Asp Leu Glu His Arg Tyr Ala
 165 170 175
 Leu Leu Gln Lys His Ile Leu Pro Ala Ile Ala Val Pro Ser Val Leu
 180 185 190
 Ala Ala Leu Asp Leu Ser Glu Glu Arg Gly Ala Arg Leu Ala Ala Glu
 195 200 205
 Leu Ala Pro Thr Gly Lys Asp
 210 215

<210> 17
 <211> 313
 <212> PRT
 <213> Streptomyces coelicolor

<400> 17
 Met Pro Glu Ala Val Val Leu Ile Asn Ser Ala Ser Asp Ala Asn Ser
 1 5 10 15
 Ile Glu Gln Thr Ala Leu Pro Val Pro Met Ala Leu Val His Arg Thr
 20 25 30
 Arg Val Gln Asp Ala Phe Pro Val Ser Trp Ile Pro Lys Gly Gly Asp
 35 40 45
 Arg Phe Ser Val Thr Ala Val Leu Pro His Asp His Pro Phe Phe Ala
 50 55 60
 Pro Val His Gly Asp Arg His Asp Pro Leu Leu Ile Ala Glu Thr Leu
 65 70 75 80
 Arg Gln Ala Ala Met Leu Val Phe His Ala Gly Tyr Gly Val Pro Val
 85 90 95
 Gly Tyr His Phe Leu Met Thr Leu Asp Tyr Thr Cys His Leu Asp His
 100 105 110
 Leu Gly Val Ser Gly Glu Val Ala Glu Leu Glu Val Glu Val Ala Cys
 115 120 125
 Ser Gln Leu Lys Phe Arg Gly Gly Gln Pro Val Gln Gly Gln Val Asp
 130 135 140
 Trp Ala Val Arg Arg Ala Gly Arg Leu Ala Ala Thr Gly Thr Ala Thr
 145 150 155 160
 Thr Arg Phe Thr Ser Pro Gln Val Tyr Arg Arg Met Arg Gly Asp Phe
 165 170 175
 Ala Thr Pro Thr Ala Ser Val Pro Gly Thr Ala Pro Val Pro Ala Ala
 180 185 190
 Arg Ala Gly Arg Thr Arg Asp Glu Asp Val Val Leu Ser Ala Ser Ser
 195 200 205

Gln Gln Asp Thr Trp Arg Leu Arg Val Asp Thr Ser His Pro Thr Leu
 210 215 220
 Phe Gln Arg Pro Asn Asp His Val Pro Gly Met Leu Leu Leu Glu Ala
 225 230 235 240
 Ala Arg Gln Ala Ala Cys Leu Val Thr Gly Pro Ala Pro Phe Val Pro
 245 250 255
 Ser Ile Gly Gly Thr Arg Phe Val Arg Tyr Ala Glu Phe Asp Ser Pro
 260 265 270
 Cys Trp Ile Gln Ala Thr Val Arg Pro Gly Pro Ala Ala Gly Leu Thr
 275 280 285
 Thr Val Arg Val Thr Gly His Gln Asp Gly Ser Leu Val Phe Leu Thr
 290 295 300
 Thr Leu Ser Gly Pro Ala Phe Ser Gly
 305 310

<210> 18
 <211> 262
 <212> PRT
 <213> Streptomyces coelicolor

<400> 18
 Met Arg Ala His Gly Thr Arg Tyr Gly Arg Pro Leu Glu Gly Lys Thr
 1 5 10 15
 Ala Leu Val Thr Gly Gly Ser Arg Gly Ile Gly Arg Gly Ile Ala Leu
 20 25 30
 Arg Leu Ala Ala Asp Gly Ala Leu Val Ala Val His Tyr Gly Ser Ser
 35 40 45
 Glu Ala Ala Ala Arg Glu Thr Val Glu Thr Ile Arg Ser Ser Gly Gly
 50 55 60
 Gln Ala Leu Ala Ile Arg Ala Glu Leu Gly Val Val Gly Asp Ala Ala
 65 70 75 80
 Ala Leu Tyr Ala Ala Phe Asp Ala Gly Met Gly Glu Phe Gly Val Pro
 85 90 95
 Pro Glu Phe Asp Ile Leu Val Asn Asn Ala Gly Val Ser Gly Ser Gly
 100 105 110
 Arg Ile Thr Glu Val Thr Glu Glu Val Phe Asp Arg Leu Val Ala Val
 115 120 125
 Asn Val Arg Ala Pro Leu Phe Leu Val Gln His Gly Leu Lys Arg Leu
 130 135 140
 Arg Asp Gly Gly Arg Ile Ile Asn Ile Ser Ser Ala Ala Thr Arg Arg
 145 150 155 160
 Ala Phe Pro Glu Ser Ile Gly Tyr Ala Met Thr Lys Gly Ala Val Asp
 165 170 175
 Thr Leu Thr Leu Ala Leu Ala Arg Gln Leu Gly Glu Arg Gly Ile Thr
 180 185 190

Val Asn Ala Val Ala Pro Gly Phe Val Glu Thr Asp Met Asn Ala Arg
 195 200 205
 Arg Arg Gln Thr Pro Glu Ala Ala Ala Ala Leu Ala Ala Tyr Ser Val
 210 215 220
 Phe Asn Arg Ile Gly Arg Pro Asp Asp Ile Ala Asp Val Val Ala Phe
 225 230 235 240
 Leu Ala Ser Asp Asp Ser Arg Trp Ile Thr Gly Gln Tyr Val Asp Ala
 245 250 255
 Thr Gly Gly Thr Ile Leu
 260

<210> 19
 <211> 4346
 <212> DNA
 <213> Streptomyces coelicolor

<400> 19
 gtogaagaag gggtaggggt cgaagcagac ggggtactcg ttcccgcca ccgggcaccg 60
 ggtctgagct cgaagacaga ccgcacagag gacgtcggct ggcggctcct gatagtgagc 120
 ggcccccgcc ggggaacaga cagcagctc catctccac tgcgcccccg gaaagtgagc 180
 accgcacccg ggaagtgacg ccacgggagg gccacgtccg cggacggatc acccctggct 240
 tcggccgaag gcttcggcgt ggtccggcgc ccagatggcg aacggcctgg cgggcgggcc 300
 cgtcacttcc cgcacggctg gacgacctg cgccttgccc ccgccccgct gccgctcggc 360
 gctctccagg aacgcgtcgg cgaagggcct cggatacttc cggagcatct gctcgcgcgc 420
 cgcctccagg ccagctcct cgaacacgag tgaccgcccc agcacctcgg agagccgcgc 480
 cgtctgctgc ctggcgggtg tggcctgggg ccgggacagc gcgtacgccc gtccctcgtg 540
 ccggggcccg gtcagtgcgc tgacggccac tccgcgatg tcgcgcggtat cgacgcaggc 600
 aacgggggac gtgcggtaca ggcgcgggac cagcgcgtcg gaccggatgg cgggcgcccc 660
 ggacagcgtg ttggacatga aggtcctggc ccgcaggaa gtcacgtcta gcccggaact 720
 gggtaacggc cgtcgtttct cgcgtgccc ccgcgtgatg aagtcgtccg cggccgggtc 780
 cccacccggc agcatggaga gcttcacag gtgcgggacg ccggcctcgc gcgcgcgcgc 840
 ccggaacagg tggctgtccg gctcgggtgg actgttcgtg acgaggaaag ccgcgccgac 900
 ccggttgagg ggcgggtcca ggcggggggc tcggcggtac tgcggcgccc agacctcgac 960
 gttcggggcg gtgacgggtc ccggttcggc ccgcggggcg aggaactctga cgggacgggt 1020
 ccggggccagc aggtggggga cctgaacggc gaccacacg gtcaagcccg tcacaagaat 1080
 cactcggggc tctctcggg cagcgaggga gggcgccctc cgaacatata tatgagggga 1140
 agggcaggat ctgccccggg ggcggaacag gcgatgttcg cggccccggg ccggtgcttc 1200
 agcggagaa cgggggggag gacaggtggt tgaggaaagc gagggtgccc tctgatgcc 1260
 ccgtgacccg acgggtgggt agccccggc ccggcccccg ccggacgtc gctggatcc 1320
 agcaggggt gtgaactcc ggtacaggga cgaacgggt gccgcggatc gacggcacga 1380
 agggcgccgg acgggtccag aggcagggc cctgcgtg cgcctcagag agcagatgc 1440
 ccggtacgtg gtcgttggg cgtgggaaga gggtaggggt actggtgtcc accgcagtc 1500
 gccaggtgt cgtgtggaa ctggcgaga ggaacagtc ctggtcgagg gtcgacgg 1560
 ccggcgccg ggggaaggg ggggtccgg ggaacagtc ggtgggagtc ggaagtcgc 1620
 ccgcacccg ccggtagat tgaggatgg tgaagcggt cgtgggagtc ccggtggag 1680
 ccgagcgtcc gggcggggag agggccagt ccacgtgtcc ctgtacgggc tgcggcgcc 1740
 ggaacttcag ctgggaacag gcaacttcca cctccagtc ccggaactcg ccgacacgc 1800
 ccaggtggtc gagggtggag ggttagtcca ggtgggcat caggaaagtg tagccacgc 1860
 gcaagcgtga ggggggtg aagacgagga tgcggcgtg acgcaaggtc tggcgatca 1920
 ccagggatc gttgtgtcc ccgtggagc ggtggaagaa cgggtggtcg tggggagga 1980
 cggggtgac ccggaacggg tccggcctc tgggtatcca gctgaacggg aacgggtcct 2040
 gcaacgtgt ccggtggaga aggcacatg gaacgggcaa tggggttgtc cagatcagat 2100
 tggcatcgga ccgagaatt atcaaaaata ctgtttggg catgggtccc cccaggaat 2160
 catgtgatc ccagctgtt tttatggcg aacgttaaga tacaactga ggggtttct 2220
 ttctatcctt ccggggggag acatgaacaa ggaagcagc atggccaaagc aggaacgggc 2280
 gacccgacg ccggaagaga tctgtgagc cggggcgag gttctggaga agcagggcta 2340
 ccaagctgc acgatacgg agatcctcaa ggtggccggg gtaaccaagg gacccctcta 2400
 cttccacttc cagtcacagg aagaactggc gctggcggtc ttccagggcc aggaacacac 2460
 acagggcgtt ccggagcaac cctccgggt gccagaaact atcgacatgg gcatgtgtt 2520
 ctgtcaccgc ttgggcacga acgtgtgtgc ccgggcgggc gtcgcgctct ccattgacca 2580

gcaggcgcac	ggtctcgatc	gcagaggacc	cttcogtgcg	tggcacgaga	cactcctgaa	2640
gctgctgaac	caggccaaag	agaacggtga	gttgctgccc	catgttggtc	ccaccgattc	2700
ggccgatctc	tacgtgggca	cgttcgcggg	gatacaggtc	gtgtccacga	cggtcagggc	2760
ctaccaggac	ctcgaaacac	gttacggcgt	gctgcagaa	cacatccctg	ccgcacatcg	2820
ggttcctctc	gtgtcgggag	cgttcgacat	ctccgagggc	cgcggagcac	gcctcggggc	2880
cgaactggca	cggacccggc	aggactgacc	cgcgaagggc	ccgcacccga	taccgacccg	2940
cctgcgcgca	ggggccgacc	ggggccggct	acgggcccgg	cggggggccc	gtaggctctg	3000
cctgcgtaac	gaagcgtggc	gggtcagaga	atcgttccgc	cgttggratc	gacgtactgg	3060
cctgtgatcc	accgtgagtc	gtcggaggcc	agaaaggcca	ccacgtccgg	gatgtcgtcg	3120
ggtctgcgca	tgcgtttgaa	cacggagttg	gcggccagtg	cggggggccc	ctcgggggtc	3180
tgcgcgcgca	gtcgttccat	gtcgtctctc	acgaaacccg	gcgcacccgc	gttgaccctg	3240
atcccccgtt	ccccagttg	cctggccagg	gcgagcgtga	gcgtgtccac	cgcacccctg	3300
gtcatcgggt	atccgatgga	ctgggggaac	gcgcgcgggg	tgcgggcaga	cagatgttgc	3360
atgatccgcc	cgcggtccgg	cagtcgtttc	agtcctgtct	ggaaccaggaa	cagcgggtgc	3420
cggacgttga	cgggcaccag	tgggtcgaag	acctcctcgg	tgacctccgt	gatccgtccc	3480
gagccgctga	cgcgcgcgtc	gttcaccagg	atgtcgaaat	cgggcgggac	tccgaactcg	3540
cccatcccgj	cgtcgaaagg	cgggtagagg	gcggccggct	cacccacgac	gcggagttcg	3600
gcgcggatgg	ccaacgcctg	tccgcgcgtg	ctccggatgg	tctcgacggc	ctctcgcgcc	3660
gcgcgcctcg	tgtgcgcgta	gtggactgac	acgagcggcc	cgtccggggc	cagccgcagg	3720
gcgataccgc	gtccgatgac	ccggcttccc	ccggtcacca	gggcggctct	gcctctcagg	3780
ggtcttccat	acctcgtccc	atgtgcacgc	atatacgccc	ccgcgcgtgg	tgagcgaccc	3840
atggcggccg	ctgggcggtt	cgaatcgacg	gtcacagcct	acctgtgacc	gcgtcagacg	3900
gggcgggagt	ggcccggttg	gacggctggg	gcagatccgg	gcggcgcgca	cggggaaccc	3960
gcgcgggtca	ggggtcaggg	gtgcgcggga	ccgcaccaggc	cggtcagggc	accgaccgga	4020
tgcaggtcgg	gcgtgcacag	cggccaccag	tctcgcgggc	ccagctccga	ctcgtacgcg	4080
taccagagcc	cggtcgggac	gagttcgagc	tggacgtggc	cgcgcgggtg	ggtgaggcgg	4140
ttgcgccagg	ggcggaaggc	ggggaggctc	gcggcgagca	tcatggggcg	ggcgcgggtc	4200
aaacggcccg	cgggcgggtc	ccagggtccc	tccaggacgt	ctagaccgcg	caaccgcgcc	4260
tgcgcgccag	cgggcagcgg	ccgcgcgcgc	tccgcgcgtg	cgcgtccggc	ggccgaggcg	4320
agcgacgcgt	agagcgcgcg	ggtacc				4346